



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/083,720
Source: OPE
Date Processed by STIC: 3/27/02

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER
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Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
 2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
 3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name,
Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name,
Sequence Information, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
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Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002



30-01-01 Comply
30-01-01 No Needed

OIFE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/083,720

DATE: 03/27/2002

TIME: 14:15:08

Input Set : A:\DX0644KBK 28FEB2002.ST25.txt

Output Set: N:\CRF3\03272002\J083720.raw

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3 <110> APPLICANT: de Waal Malefyt, Rene
4      Fickenscher, Helmut
5      Fleckenstein, Bernhard
6      Knappe, Andrea
8 <120> TITLE OF INVENTION: MAMMALIAN CYTOKINE; RELATED REAGENTS
10 <130> FILE REFERENCE: DX0644KBK
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/083,720
C--> 12 <141> CURRENT FILING DATE: 2002-02-28
12 <150> PRIOR APPLICATION NUMBER: 09/363,993
13 <151> PRIOR FILING DATE: 1999-07-29
15 <160> NUMBER OF SEQ ID NOS: 21
17 <170> SOFTWARE: PatentIn version 3.1
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 1076
21 <212> TYPE: DNA
22 <213> ORGANISM: Homo sapiens
24 <220> FEATURE:
25 <221> NAME/KEY: CDS
26 <222> LOCATION: (36)..(548)
27 <223> OTHER INFORMATION:
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31 ctgtgagtga cacacgctga gtggggtgaa gggaa atg ctg gtg aat ttc att      53
32                                     Met Leu Val Asn Phe Ile
33                                     1           5
35 ttg agg tgt ggg ttg ctg tta gtc act ctg tct ctt gcc att gcc aag      101
36 Leu Arg Cys Gly Leu Leu Leu Val Thr Leu Ser Leu Ala Ile Ala Lys
37          10           15           20
39 cac aag caa tct tcc ttc acc aaa agt tgt tac cca agg gga aca ttg      149
40 His Lys Gln Ser Ser Phe Thr Lys Ser Cys Tyr Pro Arg Gly Thr Leu
41          25           30           35
43 tcc caa gct gtt gac gct ctc tat atc aaa gca gca tgg ctc aaa gca      197
44 Ser Gln Ala Val Asp Ala Leu Tyr Ile Lys Ala Ala Trp Leu Lys Ala
45          40           45           50
47 acg att cca gaa gac cgc ata aaa aat ata cga tta tta aaa aag aaa      245
48 Thr Ile Pro Glu Asp Arg Ile Lys Asn Ile Arg Leu Leu Lys Lys Lys

55 tcc ttc ttc atg gaa gac gtt ttt gat caa cta caa ttg caa ggc tgc      341
56 Ser Phe Phe Met Glu Asp Val Phe Gly Gln Leu Gln Leu Gln Gly Cys
57          90           95           100

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60 Lys Lys Ile Arg Phe Val Glu Asp Phe His Ser Leu Arg Gln Lys Leu
61      105      110      115
63 agc cac tgt att tcc tgt gct tca tca gct aga gag atg aaa tcc att      437
64 Ser His Cys Ile Ser Cys Ala Ser Ser Ala Arg Glu Met Lys Ser Ile
65      120      125      130
67 acc agg atg aaa aga ata ttt tat agg att gga aac aaa gga atc tac      485
68 Thr Arg Met Lys Arg Ile Phe Tyr Arg Ile Gly Asn Lys Gly Ile Tyr
69 135      140      145      150
71 aaa gcc atc agt gaa ctg gat att ctt ctt tcc tgg att aaa aaa tta      533
72 Lys Ala Ile Ser Glu Leu Asp Ile Leu Leu Ser Trp Ile Lys Lys Leu
73      155      160      165
75 ttg gaa agc agt cag taaaccaaag ccaagtacat tgattttaca gttattttga      588
76 Leu Glu Ser Ser Gln
77      170
79 aatacaataa gaactgctag aaatatgttt ataacagtct atttctttta aaaacttttt      648
81 aacataatac tgacggcatg ttaggtgatt cagaatagac aagaaggatt tagtaaatta      708
83 acgttttgga tataagttgt cactaatttg cacattttct gtgttttcaa ataattgttc      768
85 cattctgaac atgttttgtc attcacaagt acattgtgtc aacttaattt aaagtatgta      828
87 acctgaatta actcgtgtaa tatttgtgtg tggagtggga tgtggggggg ggagggggaa      888
89 tgacagattt ctggaatgca atgtaatgtt actgagactt aaatagatgt tatgtatatg      948
91 attgtctgtt taagtgtttg aaaattgtta attatgccca gtgtgaactt agtacttaac      1008
93 acattttgat ttaattaaa taaattgggt ttctttctca aaaaaaaaaa aaaaaaaaaa      1068
95 aaaaaaaaaa      1076
98 <210> SEQ ID NO: 2
99 <211> LENGTH: 171
100 <212> TYPE: PRT
101 <213> ORGANISM: Homo sapiens
103 <400> SEQUENCE: 2
105 Met Leu Val Asn Phe Ile Leu Arg Cys Gly Leu Leu Leu Val Thr Leu
106 1      5      10      15
109 Ser Leu Ala Ile Ala Lys His Lys Gln Ser Ser Phe Thr Lys Ser Cys
110      20      25      30
113 Tyr Pro Arg Gly Thr Leu Ser Gln Ala Val Asp Ala Leu Tyr Ile Lys
114      35      40      45
117 Ala Ala Trp Leu Lys Ala Thr Ile Pro Glu Asp Arg Ile Lys Asn Ile
118      50      55      60
121 Arg Leu Leu Lys Lys Lys Thr Lys Lys Gln Phe Met Lys Asn Cys Gln
122 65      70      75      80
125 Phe Gln Glu Gln Leu Leu Ser Phe Phe Met Glu Asp Val Phe Gly Gln
126      85      90      95
129 Leu Gln Leu Gln Gly Cys Lys Lys Ile Arg Phe Val Glu Asp Phe His
130      100      105      110
133 Ser Leu Arg Gln Lys Leu Ser His Cys Ile Ser Cys Ala Ser Ser Ala
134      115      120      125
137 Arg Glu Met Lys Ser Ile Thr Arg Met Lys Arg Ile Phe Tyr Arg Ile

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Input Set : A:\DX0644KKBK 28FEB2002.ST25.txt

Output Set: N:\CRF3\03272002\J083720.raw

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146                               165                               170
149 <210> SEQ ID NO: 3
150 <211> LENGTH: 179
151 <212> TYPE: PRT
152 <213> ORGANISM: Equine Herpes Virus
154 <400> SEQUENCE: 3
156 Met Phe Arg Ala Ser Leu Leu Cys Cys Leu Val Leu Leu Ala Gly Val
157 1                               5                               10                               15
160 Trp Ala Asp Asn Lys Tyr Asp Ser Glu Ser Gly Asp Asp Cys Pro Thr
161                               20                               25                               30
164 Leu Pro Thr Ser Leu Pro His Met Leu His Glu Leu Arg Ala Ala Phe
165                               35                               40                               45
168 Ser Arg Val Lys Thr Phe Phe Gln Met Lys Asp Gln Leu Asp Asn Met
169                               50                               55                               60
172 Leu Leu Asp Gly Ser Leu Leu Glu Asp Phe Lys Gly Tyr Leu Gly Cys
173 65                               70                               75                               80
176 Gln Ala Leu Ser Glu Met Ile Gln Phe Tyr Leu Glu Glu Val Met Pro
177                               85                               90                               95
180 Gln Ala Glu Asn His Ser Thr Asp Gln Glu Lys Asp Lys Val Asn Ser
181                               100                              105                              110
184 Leu Gly Glu Lys Leu Lys Thr Leu Arg Val Arg Leu Arg Arg Cys His
185                               115                              120                              125
188 Arg Phe Leu Pro Cys Glu Asn Lys Ser Lys Ala Val Glu Gln Val Lys
189                               130                              135                              140
192 Ser Ala Phe Ser Lys Leu Gln Glu Lys Gly Val Tyr Lys Ala Met Ser
193 145                              150                              155                              160
196 Glu Phe Asp Ile Phe Ile Asn Tyr Ile Glu Ala Tyr Met Thr Thr Lys
197                               165                              170                              175
200 Met Lys Asn
204 <210> SEQ ID NO: 4
205 <211> LENGTH: 170
206 <212> TYPE: PRT
207 <213> ORGANISM: Epstein Barr Virus
209 <400> SEQUENCE: 4
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212 1                               5                               10                               15
215 Leu Ala Pro Glu Cys Gly Gly Thr Asp Gln Cys Asp Asn Phe Pro Gln
216                               20                               25                               30
219 Met Leu Arg Asp Leu Arg Asp Ala Phe Ser Arg Val Lys Thr Phe Phe
220                               35                               40                               45
223 Gln Thr Lys Asp Glu Val Asp Asn Leu Leu Leu Lys Glu Ser Leu Leu
224                               50                               55                               60
227 Glu Asp Phe Lys Gly Tyr Leu Gly Cys Gln Ala Leu Ser Glu Met Ile
228 65                               70                               75                               80

```

199 Ala Leu Arg Arg Leu Val Val Thr Leu Gln Cys Leu Val Leu Leu Tyr
200 Met Lys Asn

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240          115          120          125
243 Ser Lys Ala Val Glu Gln Ile Lys Asn Ala Phe Asn Lys Leu Gln Glu
244          130          135          140
247 Lys Gly Ile Tyr Lys Ala Met Ser Glu Phe Asp Ile Phe Ile Asn Tyr
248 145          150          155          160
251 Ile Glu Ala Tyr Met Thr Ile Lys Ala Arg
252          165          170
255 <210> SEQ ID NO: 5
256 <211> LENGTH: 178
257 <212> TYPE: PRT
258 <213> ORGANISM: Mus musculus
260 <400> SEQUENCE: 5
262 Met Pro Gly Ser Ala Leu Leu Cys Cys Leu Leu Leu Leu Thr Gly Met
263 1          5          10          15
266 Arg Ile Ser Arg Gly Gln Tyr Ser Arg Glu Asp Asn Asn Cys Thr His
267          20          25          30
270 Phe Pro Val Gly Gln Ser His Met Leu Leu Glu Leu Arg Thr Ala Phe
271          35          40          45
274 Ser Gln Val Lys Thr Phe Phe Gln Thr Lys Asp Gln Leu Asp Asn Ile
275          50          55          60
278 Leu Leu Thr Asp Ser Leu Met Gln Asp Phe Lys Gly Tyr Leu Gly Cys
279 65          70          75          80
282 Gln Ala Leu Ser Glu Met Ile Gln Phe Tyr Leu Val Glu Val Met Pro
283          85          90          95
286 Gln Ala Glu Lys His Gly Pro Glu Ile Lys Glu His Leu Asn Ser Leu
287          100          105          110
290 Gly Glu Lys Leu Lys Thr Leu Arg Met Arg Leu Arg Arg Cys His Arg
291          115          120          125
294 Phe Leu Pro Cys Glu Asn Lys Ser Lys Ala Val Glu Gln Val Lys Ser
295          130          135          140
298 Asp Phe Asn Lys Leu Gln Asp Gln Gly Val Tyr Lys Ala Met Asn Glu
299 145          150          155          160
302 Phe Asp Ile Phe Ile Asn Cys Ile Glu Ala Tyr Met Met Ile Lys Met
303          165          170          175
306 Lys Ser
310 <210> SEQ ID NO: 6
311 <211> LENGTH: 178
312 <212> TYPE: PRT
313 <213> ORGANISM: Homo sapiens
315 <400> SEQUENCE: 6
317 Met His Ser Ser Ala Leu Leu Cys Cys Leu Val Leu Leu Thr Gly Val
318 1          5          10          15
321 Arg Ala Ser Pro Gly Gln Gly Thr Gln Ser Glu Asn Ser Cys Thr His
322          20          25          30
325 Phe Val Ile Ile Ile Ile Ile Ile Ile Ile Ile Ile Ile Ile Ile Ile

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Input Set : A:\DX0644KBK 28FEB2002.ST25.txt

Output Set: N:\CRF3\03272002\J083720.raw

```

334 65          70          75          80
337 Gln Ala Leu Ser Glu Met Ile Gln Phe Tyr Leu Glu Glu Val Met Pro
338          85          90          95
341 Gln Ala Glu Asn Gln Asp Pro Asp Ile Lys Ala His Val Asn Ser Leu
342          100          105          110
345 Gly Glu Asn Leu Lys Thr Leu Arg Leu Arg Leu Arg Cys His Arg
346          115          120          125
349 Phe Leu Pro Cys Glu Asn Lys Ser Lys Ala Val Glu Gln Val Lys Asn
350          130          135          140
353 Ala Phe Asn Lys Leu Gln Glu Lys Gly Ile Tyr Lys Ala Met Ser Glu
354 145          150          155          160
357 Phe Asp Ile Phe Ile Asn Tyr Ile Glu Ala Tyr Met Thr Met Lys Ile
358          165          170          175
361 Arg Asn
365 <210> SEQ ID NO: 7
366 <211> LENGTH: 20
367 <212> TYPE: DNA
368 <213> ORGANISM: Synthetic - insert sequence, see over summary sheet item 16
370 <400> SEQUENCE: 7
371 tggcagcctt cctgatttct 20
374 <210> SEQ ID NO: 8
375 <211> LENGTH: 26
376 <212> TYPE: DNA
377 <213> ORGANISM: Synthetic
379 <400> SEQUENCE: 8
380 tgcactgaca tctaagttct ttagca 26
383 <210> SEQ ID NO: 9
384 <211> LENGTH: 28
385 <212> TYPE: DNA
386 <213> ORGANISM: Synthetic
388 <400> SEQUENCE: 9
389 tggcaaaact gcaccttcac acagagct 28
392 <210> SEQ ID NO: 10
393 <211> LENGTH: 21
394 <212> TYPE: DNA
395 <213> ORGANISM: Synthetic
397 <400> SEQUENCE: 10
398 gagatctccg agatgccttc a 21
401 <210> SEQ ID NO: 11
402 <211> LENGTH: 26
403 <212> TYPE: DNA
404 <213> ORGANISM: Synthetic
406 <400> SEQUENCE: 11
407 caaggactcc tttaacaaca agttgt 26

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415 <213> ORGANISM: Synthetic
416 <400> SEQUENCE: 12

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VERIFICATION SUMMARY

DATE: 03/27/2002

PATENT APPLICATION: US/10/083,720

TIME: 14:15:09

Input Set : A:\DX0644KBK 28FEB2002.ST25.txt

Output Set: N:\CRF3\03272002\J083720.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 101083,720

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0
"bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s). Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 Skipped Sequences
(OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 Skipped Sequences
(NEW RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000
- 9 Use of n's or Xaa's
(NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10 Invalid <213>
Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 Use of <220> Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses.
Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0
"bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.